

>IPI00083978.2 ACCESSION:IPI00083978 NID: Homo sapiens (Human). SIMILAR TO CD8
ANTIGEN (INDUCIBLE MEMBRANE PROTEIN R2) (C33 ANTIGEN)
(IA4) (METASTASIS SUPPRESSOR KANGAI 1) (SUPPRESSOR OF
TUMORIGENICITY- 6). IPI_human
Length = 220

Score = 451 bits (1160), Expect = e-126

Identities = 216/216 (100%), Positives = 216/216 (100%)

Query: 33 AWLLLDNRNNFLTAFDENNHFIVPISQILIGMGSSTVLFCLLGYIGIHNEIRWLLIVYAVL 92
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Sbjct: 5 AWLLLDNRNNFLTAFDENNHFIVPISQILIGMGSSTVLFCLLGYIGIHNEIRWLLIVYAVL 64

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Query: 153 CCGQHNYTDWIKNKNKENSQVPCSTKSTLRKWFCDEPLNATYLEGCENKISAWYNVNV 212
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Query: 213 LTLIGINFGLLTSEVFQVSLTVCFFKNIKNIIHAEM 248
LTLIGINFGLLTSEVFQVSLTVCFFKNIKNIIHAEM
Sbjct: 185 LTLIGINFGLLTSEVFQVSLTVCFFKNIKNIIHAEM 220

>XM_084868 ACCESSION:XM_084868 NID: gi 22059998 ref XM_084868.5 Homo sapiens
similar to CD82 antigen (Inducible membrane protein R2)
(C33 antigen) (IA4) (Metastasis suppressor Kangai 1)
(Suppressor of tumorigenicity-6) (LOC144448), mRNA
Length = 992

Score = 1465 bits (739), Expect = 0.0
Identities = 746/747 (99%), Gaps = 1/747 (0%)
Strand = Plus / Plus

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Query: 721 aagaatataatccatgcagaaatgtga 747

Sbjct: 797 aagaatataatccatgcagaaatgtga 823



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☐ 1: XM_084868. Homo sapiens simi...[gi:22059998]

Links

LOCUS LOC144448 992 bp mRNA linear PRI 01-AUG-2002
 DEFINITION Homo sapiens similar to CD82 antigen (Inducible membrane protein R2) (C33 antigen) (IA4) (Metastasis suppressor Kangai 1) (Suppressor of tumorigenicity-6) (LOC144448), mRNA.
 ACCESSION XM_084868
 VERSION XM_084868.5 GI:22059998
 KEYWORDS
 SOURCE human.
 ORGANISM Homo sapiens
 Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
 REFERENCE 1 (bases 1 to 992)
 AUTHORS NCBI Annotation Project.
 TITLE Direct Submission
 JOURNAL Submitted (31-JUL-2002) National Center for Biotechnology Information, NIH, Bethesda, MD 20894, USA
 COMMENT GENOME ANNOTATION REFSEQ: This model reference sequence was predicted from NCBI contig NT_009538 by automated computational analysis using gene prediction method: BLAST, supported by EST evidence.
 Also see:

[Documentation](#) of NCBI's Annotation Process

On Aug 1, 2002 this sequence version replaced gi:20553886.

FEATURES
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/note="Region: pfam00335, transmembrane4, Tetraspanin family"

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Revised: July 5, 2002.

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